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RAW SEQUENCE LISTING

DATE: 02/07/2002

PATENT APPLICATION: US/10/051,841

TIME: 12:53:11

Input Set : D:\AHP 98133.txt

Output Set: N:\CRF3\02072002\J051841.raw

ENTERED

SEQUENCE LISTING

4 (1) GENERAL INFORMATION:

6 (i) APPLICANT: YOUNG, KATHLEEN H.

7 RHODES, KENNETH J.

9 (ii) TITLE OF INVENTION: METHODS FOR IDENTIFYING MODULATORS OF

10 N-TYPE ION CHANNEL INACTIVATION

12 (iii) NUMBER OF SEQUENCES: 23

14 (iv) CORRESPONDENCE ADDRESS:

15 (A) ADDRESSEE: AMERICAN HOME PRODUCTS CORPORATION

16 (B) STREET: FIVE GIRALDA FARMS -- ATTN. KAY E. BRADY

17 (C) CITY: MADISON

18 (D) STATE: NEW JERSEY

19 (E) COUNTRY: US

20 (F) ZIP: 07940

22 (v) COMPUTER READABLE FORM:

23 (A) MEDIUM TYPE: CD-R

24 (B) COMPUTER: IBM PC compatible

25 (C) OPERATING SYSTEM: WINDOWS NT

26 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30

28 (vi) CURRENT APPLICATION DATA:

C--> 29 (A) APPLICATION NUMBER: US/10/051,841

C--> 30 (B) FILING DATE: 17-Jan-2002

31 (C) CLASSIFICATION:

33 (viii) ATTORNEY/AGENT INFORMATION:

34 (A) NAME: BOGLE, GAVIN T.

35 (B) REGISTRATION NUMBER:

36 (C) REFERENCE/DOCKET NUMBER: AHP 98133 C1

38 (ix) TELECOMMUNICATION INFORMATION:

39 (A) TELEPHONE: (617) 665-8079

40 (B) TELEFAX: (617) 876-5851

44 (2) INFORMATION FOR SEQ ID NO: 1:

46 (i) SEQUENCE CHARACTERISTICS:

47 (A) LENGTH: 16 amino acids

48 (B) TYPE: amino acid

49 (C) STRANDEDNESS: not relevant

W--> 50 (D) TOPOLOGY: not relevant

52 (ii) MOLECULE TYPE: peptide

54 (vi) ORIGINAL SOURCE:

55 (A) ORGANISM: Homo sapiens

59 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

61 Gln Ile Leu Gly Gln Thr Leu Lys Ala Ser Met Arg Glu Leu Gly Leu

62 1 5 10 15

65 (2) INFORMATION FOR SEQ ID NO: 2:

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67      (i) SEQUENCE CHARACTERISTICS:
68          (A) LENGTH: 16 amino acids
69          (B) TYPE: amino acid
70          (C) STRANDEDNESS: not relevant
W--> 71      (D) TOPOLOGY: not relevant
73      (ii) MOLECULE TYPE: peptide
75      (vi) ORIGINAL SOURCE:
76          (A) ORGANISM: Homo sapiens
80      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
82      Gln Ile Leu Gly His Thr Leu Arg Ala Ser Met Arg Glu Leu Gly Leu
83      1           5           10           15
86 (2) INFORMATION FOR SEQ ID NO: 3:
88      (i) SEQUENCE CHARACTERISTICS:
89          (A) LENGTH: 48 base pairs
90          (B) TYPE: nucleic acid
91          (C) STRANDEDNESS: single
92          (D) TOPOLOGY: linear
94      (ii) MOLECULE TYPE: DNA (genomic)
96      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
98      GATCCTGGGC CAGACCCTCA AAGCTAGTAT GAGAGAGCTA GGGCTGCT      48
102 (2) INFORMATION FOR SEQ ID NO: 4:
104      (i) SEQUENCE CHARACTERISTICS:
105          (A) LENGTH: 48 base pairs
106          (B) TYPE: nucleic acid
107          (C) STRANDEDNESS: single
108          (D) TOPOLOGY: linear
110      (ii) MOLECULE TYPE: DNA (genomic)
112      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
114      GATCCTGGGC CACACCCTCA GAGCCAGCAT GCGGGAAGTGGCCTTCT      48
119 (2) INFORMATION FOR SEQ ID NO: 5:
121      (i) SEQUENCE CHARACTERISTICS:
122          (A) LENGTH: 30 amino acids
123          (B) TYPE: amino acid
124          (C) STRANDEDNESS: not relevant
W--> 125      (D) TOPOLOGY: not relevant
127      (ii) MOLECULE TYPE: peptide
129      (vi) ORIGINAL SOURCE:
130          (A) ORGANISM: Homo sapiens
134      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
136      Met Gln Val Ser Ile Ala Cys Thr Glu His Asn Leu Lys Ser Arg Asn
137      1           5           10           15
139      Gly Glu Asp Arg Leu Leu Ser Lys Gln Ser Ser Thr Ala Pro
140      20           25           30
142 (2) INFORMATION FOR SEQ ID NO: 6:
144      (i) SEQUENCE CHARACTERISTICS:
145          (A) LENGTH: 30 amino acids
146          (B) TYPE: amino acid
147          (C) STRANDEDNESS: not relevant
W--> 148      (D) TOPOLOGY: not relevant

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150      (ii) MOLECULE TYPE: peptide
152      (vi) ORIGINAL SOURCE:
153          (A) ORGANISM: Homo sapiens
157      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
159      Met Glu Val Ala Met Val Ser Ala Glu Ser Ser Gly Cys Asn Ser His
160      1              5              10              15
162      Met Pro Tyr Gly Tyr Ala Ala Gln Ala Arg Ala Arg Glu Arg
163      20              25              30
166 (2) INFORMATION FOR SEQ ID NO: 7:
168      (i) SEQUENCE CHARACTERISTICS:
169          (A) LENGTH: 90 base pairs
170          (B) TYPE: nucleic acid
171          (C) STRANDEDNESS: single
172          (D) TOPOLOGY: linear
174      (ii) MOLECULE TYPE: DNA (genomic)
176      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
178      ATGCAAGTCT CCATAGCCTG CACAGAGCAC AATTTGAAGA GTCGGAATGG TGAGGACCGA      60
180      CTTCTGAGCA AGCAGAGCTC CACGCCCCC      90
185 (2) INFORMATION FOR SEQ ID NO: 8:
187      (i) SEQUENCE CHARACTERISTICS:
188          (A) LENGTH: 90 base pairs
189          (B) TYPE: nucleic acid
190          (C) STRANDEDNESS: single
191          (D) TOPOLOGY: linear
193      (ii) MOLECULE TYPE: DNA (genomic)
195      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
197      ATGGAGGTTG CAATGGTGAG TCGGAGAGC TCAGGGTGCA ACAGTCACAT GCCTTATGGT      60
199      TATGCTGCCC AGGCCCGGGC CCGGGAGCGG      90
202 (2) INFORMATION FOR SEQ ID NO: 9:
204      (i) SEQUENCE CHARACTERISTICS:
205          (A) LENGTH: 59 base pairs
206          (B) TYPE: nucleic acid
207          (C) STRANDEDNESS: single
208          (D) TOPOLOGY: linear
210      (ii) MOLECULE TYPE: DNA (genomic)
213      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
215      CATGGAGCTC TTCATCGGGG TCATCCTGTT TTCTAGTGCA GTGTACTTTG CCGAGTAAG      59
218 (2) INFORMATION FOR SEQ ID NO: 10:
220      (i) SEQUENCE CHARACTERISTICS:
221          (A) LENGTH: 59 base pairs
222          (B) TYPE: nucleic acid
223          (C) STRANDEDNESS: single
224          (D) TOPOLOGY: linear
226      (ii) MOLECULE TYPE: DNA (genomic)
228      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
230      GATCCTTACT CGCAAAGTA CACTGCACTA GAAACAGGA TGACCCCGAT GAAGAGCTC      59
233 (2) INFORMATION FOR SEQ ID NO: 11:
235      (i) SEQUENCE CHARACTERISTICS:
236          (A) LENGTH: 55 base pairs

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237         (B) TYPE: nucleic acid
238         (C) STRANDEDNESS: single
239         (D) TOPOLOGY: linear
241     (ii) MOLECULE TYPE: DNA (genomic)
245     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
247 CCGAATTCGA CATATGAAAA TGCAAGTCTC CATAGCCTGC ACAGAGCACA ATTTG
250 (2) INFORMATION FOR SEQ ID NO: 12:
252     (i) SEQUENCE CHARACTERISTICS:
253         (A) LENGTH: 42 base pairs
254         (B) TYPE: nucleic acid
255         (C) STRANDEDNESS: single
256         (D) TOPOLOGY: linear
258     (ii) MOLECULE TYPE: DNA (genomic)
260     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
262 ACGGATCCCC GAATTCCATT ATGATCTATA GTCCTTCTTG CT
265 (2) INFORMATION FOR SEQ ID NO: 13:
267     (i) SEQUENCE CHARACTERISTICS:
268         (A) LENGTH: 45 base pairs
269         (B) TYPE: nucleic acid
270         (C) STRANDEDNESS: single
271         (D) TOPOLOGY: linear
273     (ii) MOLECULE TYPE: DNA (genomic)
275     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
277 ACGGATCCCC GAATTCCATT AATCTGAAAT TTGACCTCCA AATGT
280 (2) INFORMATION FOR SEQ ID NO: 14:
282     (i) SEQUENCE CHARACTERISTICS:
283         (A) LENGTH: 59 base pairs
284         (B) TYPE: nucleic acid
285         (C) STRANDEDNESS: single
286         (D) TOPOLOGY: linear
288     (ii) MOLECULE TYPE: DNA (genomic)
290     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
292 CATGGAGCAG ATCCTGGGCC ACACCCTGAG AGCCAGCATG CGGGAAGTGG GCCTTTAAG
295 (2) INFORMATION FOR SEQ ID NO: 15:
297     (i) SEQUENCE CHARACTERISTICS:
298         (A) LENGTH: 59 base pairs
299         (B) TYPE: nucleic acid
300         (C) STRANDEDNESS: single
301         (D) TOPOLOGY: linear
303     (ii) MOLECULE TYPE: DNA (genomic)
305     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
307 GATCCTTAAA GGCCAGTTC CGGCATGCTG GCTCTGAGGG TGTGGCCCAG GATCTGCTC
310 (2) INFORMATION FOR SEQ ID NO: 16:
312     (i) SEQUENCE CHARACTERISTICS:
313         (A) LENGTH: 60 base pairs
314         (B) TYPE: nucleic acid
315         (C) STRANDEDNESS: single
316         (D) TOPOLOGY: linear
318     (ii) MOLECULE TYPE: DNA (genomic)

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320      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
322 CGAATTCATA TGCGGATCCG TAGAATGGAG GTTGCAATGG TGAGTGC GGA GAGCTCAGGG      60
325 (2) INFORMATION FOR SEQ ID NO: 17:
327      (i) SEQUENCE CHARACTERISTICS:
328          (A) LENGTH: 41 base pairs
329          (B) TYPE: nucleic acid
330          (C) STRANDEDNESS: single
331          (D) TOPOLOGY: linear
333      (ii) MOLECULE TYPE: DNA (genomic)
335      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
337 GGTCGACGAA TCGTTACCT TGCAGGATCG GAGCTCTCGT G      41
340 (2) INFORMATION FOR SEQ ID NO: 18:
342      (i) SEQUENCE CHARACTERISTICS:
343          (A) LENGTH: 59 base pairs
344          (B) TYPE: nucleic acid
345          (C) STRANDEDNESS: single
346          (D) TOPOLOGY: linear
348      (ii) MOLECULE TYPE: DNA (genomic)
350      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
352 CATGGAGCAG ATCCTGGGCC ACACCCTCAG AGCCAGCATG CGGCAACTGG GCCTTTAAG      59
355 (2) INFORMATION FOR SEQ ID NO: 19:
357      (i) SEQUENCE CHARACTERISTICS:
358          (A) LENGTH: 59 base pairs
359          (B) TYPE: nucleic acid
360          (C) STRANDEDNESS: single
361          (D) TOPOLOGY: linear
363      (ii) MOLECULE TYPE: DNA (genomic)
365      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
367 GATCCTTAAA GGCCAGTTG CCGCATGCTG GCTCTGAGGG TGTGGCCCAG GATCTGCTC      59
370 (2) INFORMATION FOR SEQ ID NO: 20:
372      (i) SEQUENCE CHARACTERISTICS:
373          (A) LENGTH: 59 base pairs
374          (B) TYPE: nucleic acid
375          (C) STRANDEDNESS: single
376          (D) TOPOLOGY: linear
378      (ii) MOLECULE TYPE: DNA (genomic)
380      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
382 AATTCCTCTT CATCGGGGTC ATCCTGTTT CTAGTGCAGT GTACTTTGCC GAGTAAGCC      59
385 (2) INFORMATION FOR SEQ ID NO: 21:
387      (i) SEQUENCE CHARACTERISTICS:
388          (A) LENGTH: 59 base pairs
389          (B) TYPE: nucleic acid
390          (C) STRANDEDNESS: single
391          (D) TOPOLOGY: linear
393      (ii) MOLECULE TYPE: DNA (genomic)
395      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
397 TCGAGGCTTA CTCGGCAAAG TACACTGCAC TAGAAAACAG GATGACCCCG ATGAAGAGG      59
400 (2) INFORMATION FOR SEQ ID NO: 22:
402      (i) SEQUENCE CHARACTERISTICS:

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/051,841

DATE: 02/07/2002

TIME: 12:53:12

Input Set : D:\AHP 98133.txt

Output Set: N:\CRF3\02072002\J051841.raw

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:50 M:246 W: Invalid value of Alpha Sequence Header Field, [TOPOLOGY:], SeqNo=1
L:71 M:246 W: Invalid value of Alpha Sequence Header Field, [TOPOLOGY:], SeqNo=2
L:125 M:246 W: Invalid value of Alpha Sequence Header Field, [TOPOLOGY:], SeqNo=5
L:148 M:246 W: Invalid value of Alpha Sequence Header Field, [TOPOLOGY:], SeqNo=6